



1600

RAW SEQUENCE LISTING

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:21:50

Input Set : N:\efs\09865018\09865018A\PTOMS.txt

Output Set: N:\CRF4\04232003\I865018B.raw

4 <110> APPLICANT: Massague et al.
 6 <120> TITLE OF INVENTION: ISOLATED p27 PROTEIN AND METHODS FOR ITS PRODUCTION AND USE
 8 <130> FILE REFERENCE: GPC1-P03-079
 10 <140> CURRENT APPLICATION NUMBER: 09/865018B
 11 <141> CURRENT FILING DATE: 2001-05-24
 13 <150> PRIOR APPLICATION NUMBER: 08/854039
 14 <151> PRIOR FILING DATE: 1997-05-09
 16 <160> NUMBER OF SEQ ID NOS: 27
 17 <170> SOFTWARE: Patentix version 3.1
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 597
 22 <212> TYPE: DNA
 23 <213> ORGANISM: Homo sapiens
 24 <210> FEATURES:
 26 <211> NAME/KEY: CDS
 27 <212> LOCATION: (1)..(597)
 29 <213> OTHER INFORMATION:

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52 Met Ser Asn Val Arg Val Ser Asn Gly Ser Pro Ser Leu Glu Arg Met	
53 1 5 10 15	
54 aac gcc agg cag gcc gag cac cca aag ccc tgg gcc tgc agg aac ctg	96
55 Asp Ala Arg Gln Ala Glu His Pro Lys Pro Ser Ala Cys Arg Asn Leu	
56 20 25 30	
57 ttc ggt cgg gng gac cac gaa gag tta acc cgg gac ttg gag aag cac	144
58 Phe Gly Pro Val Asp His Glu Gln Leu Thr Arg Asp Leu Glu Lys His	
59 35 40 45	
60 tgc aga gac atg caa gag gcc agc cag cgc aag tgg aat ttc gat ttt	192
61 Cys Arg Asp Met Gln Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe	
62 50 55 60	
63 caa aat cac aaa ccc cta gag gcc aag tac gag tgg caa gag gtg gag	240
64 Gln Asn His Lys Pro Leu Glu Gly Lys Tyr Glu Trp Gln Glu Val Glu	
65 65 70 75 80	
66 aag gcc agc ttg ccc gag ttc tac tac aga ccc ccg cgg ccc ccc aaa	288
67 Lys Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys	
68 85 90 95	
69 ggt gcc tgc aag gtg ccg gcc caa gag agc caa gat gtc agc cgg agc	336
70 Gly Ala Cys Lys Val Pro Ala Gln Glu Ser Gln Asp Val Ser Gly Ser	
71 100 105 110	
72 cgc ccg gcc gcc cct tta att gga gct ccg gct aac tct gag gac acc	384
73 Arg Pro Ala Ala Pro Leu Ile Gly Ala Pro Ala Asn Ser Glu Asp Thr	
74 115 120 125	
75 cat ttg gtg gac cca aag act gat ccg tgg gac acc cag acc gga tta	432

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Output Set: N:\CRF4\04232003\I865018B.raw

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64 His Leu Val Asp Pro Lys Thr Asp Pro Ser Asp Ser Gln Thr Gly Leu
65      130      135      140
67 ggc gag caa ttc gca gga ata agg aag oga cct gca acc gac gat tct      480
68 Ala Glu Gln Cys Ala Gly Ile Arg Lys Arg Pro Ala Thr Asp Asp Ser
69 145      150      155      160
71 tct act caa aac aat aga gcc aac aga aca gaa gaa aat gtt tca gat      528
72 Ser Thr Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
73      165      170      175
75 ggt tcc cca aat gac ggt tct gtg gag cag aag ccc aag aag cct ggc      576
76 Gly Ser Pro Asn Ala Gly Ser Val Glu Gln Thr Pro Lys Lys Pro Gly
77      180      185      190
79 ctc aga aga cgt cca aag taa      597
80 Leu Arg Arg Arg Gln Thr
81      195
83 <210> SEQ ID NO: 2
84 <211> LENGTH: 198
85 <212> TYPE: PRT
86 <213> ORGANISM: Homo sapiens
87 <400> SEQUENCE: 2
89 Met Ser Asn Val Arg Val Ser Asn Gly Ser Pro Ser Leu Glu Arg Met
90      5      10      15
94 Asp Ala Arg Gln Ala Glu His Pro Lys Pro Ser Ala Cys Arg Asn Leu
95      20      25      30
98 Phe Gly Pro Val Asp His Glu Glu Leu Thr Arg Asp Leu Glu Lys His
99      35      40      45
102 Cys Arg Asp Met Glu Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
103      50      55      60
106 Gln Asn His Lys Pro Leu Glu Gly Lys Tyr Glu Trp Gln Glu Val Glu
107 65      70      75      80
110 Lys Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
111      85      90      95
114 Gly Ala Cys Lys Val Pro Ala Gln Glu Ser Gln Asp Val Ser Gly Ser
115      100      105      110
118 Arg Pro Ala Ala Pro Leu Ile Gly Ala Pro Ala Asn Ser Glu Asp Thr
119      115      120      125
122 His Leu Val Asp Pro Lys Thr Asp Pro Ser Asp Ser Gln Thr Gly Leu
123      130      135      140
126 Ala Glu Gln Cys Ala Gly Ile Arg Lys Arg Pro Ala Thr Asp Asp Ser
127 145      150      155      160
130 Ser Thr Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
131      165      170      175
134 Gly Ser Pro Asn Ala Gly Ser Val Glu Gln Thr Pro Lys Lys Pro Gly
135      180      185      190
138 Leu Arg Arg Arg Gln Thr
139      195
141 <210> SEQ ID NO: 3
142 <211> LENGTH: 594
143 <212> TYPE: DNA
144 <213> ORGANISM: Mus musculus

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PATENT APPLICATION: US/09/865,018B

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Input Set : N:\efs\09865018\09865018A\PTOMS.txt

Output Set: N:\CRF4\04232003\I865018B.raw

146 <100> FEATURE:

147 <101> NAME/KEY: CDS

148 <102> LOCATION: (1)..(594)

149 <103> OTHER INFORMATION:

W--> 151 <400> 3

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153 1      5      10      15
154 gac gcc aga caa ggg gat cac ccc aag cct tcc gcc tgc aga aat ctg      96
155 Asp Ala Arg Gln Ala Asp His Pro Lys Pro Ser Ala Cys Arg Asn Leu
156 20      25      30
157 ttc ggc cgg gtc aat cat gaa gaa cta acc cgg gac ttg gag aag cac      144
158 Phe Gly Pro Val Asn His Glu Glu Leu Thr Arg Asp Leu Glu Lys His
159 35      40      45
160 tcc cgg gat atg gaa gaa ggg agt cag cgc aag tgg aat ttc gac ttt      192
161 Cys Arg Asp Met Glu Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
162 50      55      60
163 cag aat cat aag ccc ctg gag ggc aga tac gaa tgg cag gag gtg gag      240
164 Gln Asn His Lys Pro Leu Glu Gly Arg Tyr Glu Trp Gln Glu Val Glu
165 65      70      75      80
166 aag gcc agc ttc ccc gag ttc tac tac agg ccc cgg cgc ccc ccc aag      288
167 Arg Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
168 85      90      95
169 gac gcc tgc aag gtg ctg ggc cag gag agc cag gat gtc agc ggg agc      336
170 Ser Ala Cys Lys Val Leu Ala Gln Glu Ser Gln Asp Val Ser Gly Ser
171 100      105      110
172 cgc cag gag gtg cct tta att ggg tot cag gca aac tot gag gac cgg      384
173 Arg Gln Ala Val Pro Leu Ile Gly Ser Gln Ala Asn Ser Glu Asp Arg
174 115      120      125
175 cat ttg gtg gac caa atg cct gac tgg tca gac aat cag gct ggg tta      432
176 His Leu Val Asp Gln Met Pro Asp Ser Ser Asp Asn Gln Ala Gly Leu
177 130      135      140
178 gag cag cag tot cca ggg atg agg aag cga cct gct gca gaa gat tot      480
179 Ala Glu Gln Cys Pro Gly Met Arg Lys Arg Pro Ala Ala Glu Asp Ser
180 145      150      155      160
181 tct tgg caa aac aaa agg gcc aac aga aca gaa gaa aat gtt tca gac      528
182 Ser Ser Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
183 165      170      175
184 gat tcc cgg aac gct ggc act gtg gag cag acg ccc aag aag ccc ggc      576
185 Gly Ser Pro Asn Ala Gly Thr Val Glu Gln Thr Pro Lys Lys Pro Gly
186 180      185      190
187 att cga cgc cag acg taa
188 Leu Arg Arg Gln Thr
189 195

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204 <100> SEQ ID NO: 4

205 <101> LENGTH: 195

206 <102> TYPE: PRT

207 <103> ORGANISM: Mus musculus

209 <400> SEQUENCE: 4

RAW SEQUENCE LISTING

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:21:50

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Output Set: N:\CRF4\04232003\I865018B.raw

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211 Met Ser Asn Val Arg Val Ser Asn Gly Ser Pro Ser Leu Glu Arg Met
212 1 10 15
215 Asp Ala Arg Gln Ala Asp His Pro Lys Pro Ser Ala Cys Arg Asn Leu
216 20 25 30
219 Phe Gly Pro Val Asn His Glu Gln Leu Thr Arg Asp Leu Glu Lys His
220 35 40 45
223 Cys Arg Asp Met Glu Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
224 50 55 60
227 Gln Asn His Lys Pro Leu Glu Gly Arg Tyr Gln Trp Gln Glu Val Glu
228 65 70 75 80
231 Arg Gly Ser Leu Pro Gln Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
232 85 90 95
235 Ser Ala Cys Lys Val Leu Ala Gln Glu Ser Gln Asp Val Ser Gly Ser
236 100 105 110
239 Arg Gln Ala Val Pro Leu Ile Gly Ser Gln Ala Asn Ser Glu Asp Arg
240 115 120 125
243 His Leu Val Asp Gln Met Pro Asp Ser Ser Asp Asn Gln Ala Gly Leu
244 130 135 140
247 Ala Gln Gln Cys Pro Gly Met Arg Lys Arg Pro Ala Ala Glu Asp Ser
248 145 150 155 160
251 Ser Ser Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
252 165 170 175
255 Gly Ser Pro Asn Ala Gly Thr Val Glu Gln Thr Pro Lys Lys Pro Gly
256 180 185 190
259 Leu Arg Arg Gln Thr
260 195

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261 <210> SEQ ID NO: 5

262 <211> LENGTH: 534

263 <212> TYPE: DNA

264 <213> ORGANISM: Mustela vison

265 <214> FEATURE:

266 <215> NAME/KEY: CDS

267 <216> LOCATION: (1)..(534)

268 <217> OTHER INFORMATION:

W--> 272 <400> 5

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275 1 10 15
278 gac gcc aga cag cgc gag tac ccc aag ccc tcc gcc tgc aga aac ctc 96
279 Asp Ala Arg Gln Ala Glu Tyr Pro Lys Pro Ser Ala Cys Arg Asn Leu
280 20 25 30
283 ttc ggc cgg gtc aac cac gaa gag ctg acc cgg gac ttg gag aag cac 144
284 Phe Gly Pro Val Asn His Glu Gln Leu Thr Arg Asp Leu Glu Lys His
285 35 40 45
288 ccc aga gac atg gaa gag gca agc cag cgc aag tgg aat ttt gat ttc 192
289 Arg Arg Asp Met Glu Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
290 50 55 60
293 cag aat cac aag ccc ctg gag ggc aaa tac gag tgg cag gag gtg gag 240
294 Gln Asn His Lys Pro Leu Glu Gly Lys Tyr Glu Trp Gln Glu Val Glu

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RAW SEQUENCE LISTING

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:21:50

Input Set : N:\efs\09865018\09865018A\PTOMS.txt

Output Set: N:\CRF4\04232003\I865018B.raw

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291 65          70          75          80
293 aug ggc agc ttg cgg gag ttc tac tac aga ccc cgg cgg cca ccc aac 288
294 Lys Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
344          85          90          95
397 ggc ggc tgc atg ttg cgg cgg cag gag agc cag gac gtc agc ggc aac 336
398 Gly Ala Cys Lys Val Pro Ala Gln Glu Ser Gln Asp Val Ser Gly Thr
399          100          105          110
401 cgg cag gcc gtc cct tta atg ggg tct cag gca aac tca gag gat aca 384
402 Arg Gln Ala Val Pro Leu Met Gly Ser Gln Ala Asn Ser Glu Asp Thr
403          115          120          125
405 cag ttg gta gac aac aag act gac aag cgg gac aac cag gct ggc tta 432
406 His Leu Val Asp Gln Lys Thr Asp Thr Ala Asp Asn Gln Ala Gly Leu
407          130          135          140
409 gag aag cag ttc act ggg atc agc aag cga cgg gcc aca gac gat ttc 480
410 Ala Glu Gln Cys Thr Gly Ile Arg Lys Arg Pro Ala Thr Asp Asp Ser
411 141          150          155          160
413 cct cct aac aac aag aga gcc aac aga aca gac gac aat gtc tca gac 528
414 Ser Pro Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
415          165          170          175
417 ggt tcc
418 Gly Ser
421 4210 - SEQ ID NO: 6
422 4211 - LENGTH: 177
423 4212 - TYPE: PRT
424 4213 - ORGANISM: Mustela vison
425 4240 - SEQUENCE: 6
426 Met Ser Asn Val Arg Val Ser Asn Gly Ser Pro Ser Leu Glu Arg Met
427 1          5          10          15
428 Asp Ala Arg Gln Ala Glu Tyr Pro Lys Pro Ser Ala Cys Arg Asn Leu
429          20          25          30
430 Phe Gly Pro Val Asn His Glu Glu Leu Thr Arg Asp Leu Glu Lys His
431          35          40          45
432 Arg Arg Asp Met Glu Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
433          50          55          60
434 Gln Asn His Lys Pro Leu Glu Gly Lys Tyr Glu Trp Gln Glu Val Glu
435 65          70          75          80
438 Lys Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
439          85          90          95
442 Gly Ala Cys Lys Val Pro Ala Gln Glu Ser Gln Asp Val Ser Gly Thr
443          100          105          110
446 Arg Gln Ala Val Pro Leu Met Gly Ser Gln Ala Asn Ser Glu Asp Thr
447          115          120          125
450 His Leu Val Asp Gln Lys Thr Asp Thr Ala Asp Asn Gln Ala Gly Leu
451          130          135          140
454 Ala Glu Gln Cys Thr Gly Ile Arg Lys Arg Pro Ala Thr Asp Asp Ser
455 145          150          155          160
458 Ser Pro Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
459          165          170          175
472 Gly Ser

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/865,018B

DATE: 04/23/2003
TIME: 07:21:51

Input Set : N:\efs\09865018\09865018A\PTOMS.txt
Output Set: N:\CRF4\04232003\I865018B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:12; N Pos. 3,9
Seq#:13; N Pos. 1,13,16,22
Seq#:14; N Pos. 6,9,12,15
Seq#:15; N Pos. 6,9,12
Seq#:16; N Pos. 1,7,10,13,16
Seq#:17; N Pos. 1,10,13,16
Seq#:18; N Pos. 3
Seq#:19; N Pos. 15
Seq#:21; N Pos. 1

VERIFICATION SUMMARY

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:21:51

Input Set : N:\efs\09865018\09865018A\PTOMS.txt

Output Set: N:\CRF4\04232003\I865018B.raw

L:30 M:258 W: Mandatory Feature missing, <203> Blank for SEQ#:1,Line#:28
L:151 M:258 W: Mandatory Feature missing, <220> Blank for SEQ#:3,Line#:149
L:272 M:258 W: Mandatory Feature missing, <220> Blank for SEQ#:5,Line#:270
L:442 M:341 W: (46) "r" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:459 M:341 W: (46) "r" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:466 M:283 W: Missing Blank Line separator, <220> field identifier
L:474 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:536 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:554 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0



Does Not Comply
Corrected Diskette Needed

1600

RAW SEQUENCE LISTING

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:17:28

Input Set : N:\efs\09865018\09865018A\GPCI-P08-079SubstituteSequence.txt

Output Set: N:\CRF4\04232003\I865018B.raw

4 <110> APPLICANT: Massague et al.
 6 <120> TITLE OF INVENTION: ISOLATED p27 PROTEIN AND METHODS FOR ITS PRODUCTION AND USE
 8 <130> FILE REFERENCE: GPCI-P08-079
 10 <140> CURRENT APPLICATION NUMBER: 09/865018B
 11 <141> CURRENT FILING DATE: 2001-05-24
 13 <150> PRIOR APPLICATION NUMBER: 08/854039
 14 <151> PRIOR FILING DATE: 1997-05-09
 16 <160> NUMBER OF SEQ ID NOS: 27
 18 <170> SOFTWARE: PatentIn version 3.1

ERRORED SEQUENCES

643 <110> SEQ ID NO: 25
 644 <111> LENGTH: 6
 645 <112> TYPE: PRT
 646 <113> ORGANISM: Homo sapiens
 648 <140> SEQUENCE: 25
 650 Leu Phe Gly Pro Val Asp

E--> 651 1 5

Realign amino acid numbering

VERIFICATION SUMMARY

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:17:29

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Output Set: N:\CRF4\04232003\I865018B.raw

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L:151 M:158 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:149
L:272 M:158 W: Mandatory Feature missing, <223> Blank for SEQ#:5,Line#:270
L:440 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:459 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:465 M:183 W: Missing Blank Line separator, <220> field identifier
L:474 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:520 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:538 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:554 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:681 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:25

ENTERED

Changed a file from non-ASCII to ASCII

Changed the margins in cases where the sequence text was "wrapped" down to the next line.

Edited a format error in the Current Application Data section, specifically:

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

Added the mandatory heading and subheadings for "Current Application Data".

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

Inserted or corrected a nucleic number at the end of a nucleic line: SEQ ID NO's edited:

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included:

Deleted extra, invalid, headings used by an applicant, specifically:

Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;
☐ page numbers throughout text; ☐ other invalid text, such as _____.

Inserted mandatory headings, specifically:

Corrected an obvious error in the response, specifically:

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically:

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

Other:

Realigned amino acid numbering in

Set 10 T25

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form. 3/1/95

3/1/95